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Fig. 1

EGP-2 promoter sequences

B₂II

-3967. **AGATCTGAA TAGAGGGAA** TTTGGCTGCA T AGTGGTTAAG GACTTTTACT CTTCATCTCA TATAAAGGAC TTTTGTTC
-3887. TACTCATCTA TTACTTATGG GATAACAAA ATTTCAGAA CTGGTAGCTC AATTATATAT ATATATATAT ATATATATAT
-3807. ATATATATAT ATATATATAT ATATATTTT TTTTTTTT TTTTAGACAG AGTTTGGCTC TTTGGGCCA GGCTGGAGTG
-3727. CAATGGCATG ATCTTCGGTC ACCACAACT CCGCTCTCTG GTTCAAGTG ATTCTCTCTGC CTCAGGCTCC CAAGTATCTG
-3647. GAATTACAGG CATGTGCCAC CATGTGCCAC TAATTCTTAAT ATTTCAGAA GAGACAGGT TTCACCCAGGT TGCCCAAGGT
-3567. GCTCTCAABC TCTGTGACCTC AAGTGATCCA CCCGGTTGG CTCCTCCAAAG TGTGGGATT ACAGGGTGA GCCACCATGC
-3487. CTAGGCTGAA ATATTAATAAATGTGCTTA AATATGGCAC TAGAACTACA AAAGATTCAC AATTAACAA TAAACAGGT
-3407. AATTTGAGC AAGAAATGAC AAATTGAGAA GGT₁TTAATG AGGTACTAA AATAAACAAATA CCGGGCCG (GTGCA GTGGCTCA p39⁸
-3327. TGCCTGTAAT CCCAGGACTT TGGGAGCTG AGGCTGGC ATCACCTGAG GTCAAGGAGT CAAGACCC CTTGGCCAACCG
-3247. TAGTGAACC CGGCTCTAC TAAAATACA AAATTAAGCC GGGGAGGTG GCAGGGCCT. GAAATCACAG CTACTCGGGA
-3167. GGCTGAGACA GGAGAATTGC TTGZACCCAG GAGGTGAGG TTGCACTGAG CTGAGAACAC GCATTTGTAC TCCAGCCTGG
-3087. GTAAACAGAT TGAACCTTA TCTTAAGAA AAAAAGG CGGACACGGT GGCTTGACCC TGTATCTCA GCACCTTGGG
-3007. AGGCCGAGGC AAGGAGATCA CAAAGTCAGG AGATCAAGAC CATCTGGCC CAGCTTATTC AGGAGGTGA GGCAAGGAA TTGCTTGAAC
-2927. TACAAATT AGCCGGCTCT GGTGGGGCC GCCTGTAATC CAAAGTCAGG TCACTGCCACT GCAGTGCAGC TTGGGTGACA GAGGAAGGCC CCATCTCAA
-2847. CCAAGAGGTG GAGGTGCGAG TCCGCCAATC TCACTGCCACT GCAGTGCAGC TTGGGTGACA GAGGAAGGCC CCATCTCAA
-2767. AAAAAGAAA MAGAAAAT ACCCTGGATC AGGGGGTGT GGTGGCTCA GCCTGTAATC CCAGACTTTGGGGCTGA p39⁸
-2687. GGCTGGCAGA TCACTGTAGG TCAGGGCTC AAGGCCAGCC TGACCAACAT GGAGAAACCC CATTCTACT AAAAATCAA
-2607. AAAATAGCC GGACGTGGTG GCACATGCTT GAAATCCAG CTACTCAGG GGCTGAGGA GGGAATTGC CTGAATCCGG
-2547. GAGGGGGAGG TTGTTGGAG GTGAGATGAT GCAATTGCACT TCCAGGCTG GCAACAAAGG CAAACTCTG CCTCAAAA
-2447. AGAAGAGAA AAAAAGAGA AAGAAAGAA AAAATACCCCT GGATGTATACAA TGAGATACAA TGAGTCTGG ATTAGTCTGG
-2367. TATTGTCA TTATTAAT ACTAAATTCA CTTTATTAACTAAATAAATAGTGTCA CAGTTATAAG p39⁸
-2287. AAGATGAAGT TCTCCCGATT AGGTAAACAG ATTAGACCT CAGAATGQA CATTGTCGA ATAAAGCCAC ATAAACAGT
-2207. TAGTTATTC TTGGGAAAG TATATGAT TTGGAGAAAG GAAACATC CAAATTCA CAGACAAACAA
-2127. AAATCTGGTT AACCTGTCTC TGAATTGTTA GAACTATTCTTC TTTTTTTTTG TTGTTGTT TTTTTTTTTT GAGACGGAGT p39⁸
-2047. TTGCTCTTG TTGCCAGGC TGGAGTGCAA TGGGAAATG S₁ TGGGTCACT GCAACCTCTG CCTCCAGGT
-1967. CTCCCTGCCCTC AGTCTCTGAA GTAGCTGGAA TTACAGGGCA GCCTGGCTAA CTTCTGTAT TTTTAGTAGA
-1887. GACGGGGTTT CACCATGTTT GCCAGGGCTGG TCTGAACTC CTGACCTTAG GTGATCCCTC S₁ GGTCTGGCC TCCCAAGGTG
-1807. CTGAAGATTAC AGGGATGAGC CACCGTACCT GGCTTAATA CCTTATTCA TATACCAAGT GAAATTAAA TTATACAAA

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Fig. 1, contd.

-1727. CAAATTAG AGGTACTTAG AACAGGATG AATTCACAT TAATCAACTT GCCGGCACTT CAACAGAATA CAACATAGAA

-1647. ATGATTGTT TAATTAAC ATAAAGCTTG ATTGACATA TACTGTAGA ATTAATCAA ACTTAGCTGA ATCTTAAAT

-1567. TGGTTTTTA CCGTCTCTT TTTTTTTA TTTTTTATT TTTTGGATG GAGTCTTGT CTGTTGCCAG ACTGGAGTGC

-1487. AGGGGTTGG TCTCGGTCA CGCGAACCTC CGACTCTCTG GTTCAGGGA TTCTCCTGC TCAGCCCT GAGTAGCTGG

-1407. GTTACAGGT GCCTGCCACC ACACCTGGT ACTTTTGTAA TTTTAGTTG AGATGGGTT CACCATGTTG GCCAGGATGG

-1327. TCTCGAACTC CTGACCTCGG ATCTGCCAAC CTGTCCTCGG AGCAAGGTGC TGGATTACA AGCATGAGCC ACCGTGCCCA

-1247. GCCTCCTTC CTCCTTAA CTCCTTACTT TATGTTCT TTAGTTCTT AAAGCTTAAAGTTAGG TTACATGAT

-1167. ATTACAGGTA ACAAAATA ACATTAATA ACACTAAATA GTATATATAT GAAGTATTAA TAATTATTT AATATTGAA

-1087. TAAATAGTG TGTGTGATT TGAATTCACTC TGACCGAAA TGATTAAC TGCTTTCTT CTATTCCTT ATATTTCTT p39^{st-2}

-1017. TCCGAGGTT CATCAACATT TTGGTTCTT AAATAGTACCC AAAACCGAA ATCATCTGG TTCTCAGTAT TTGGCTCTAT

-937. GGGAACTCTT TTCTTTCTT CTCTTTTTT TTTTTTGTAA GACGGAGTCT TGCTCCTGTC GCCCAGGCTG GAGGTAAATG

-857. GCAAGATCTC TGCTCCTGC AACCTCAGCC TCCCAGTAG CTGGGATTAC AGGCATGCGC CACCAAGCCC GGCTAAATT

-777. GTATCTTTA GTAGAGACGG CGCTCTTCAG TGTGGTCTG GCTGGTCTG AACTCAAC CTCAGGTGAT CGGCCGCCCT

-697. CGGCCCTCCC AAAGTGTAGG ATTACAGGG TGACCCACCG CGCTCAGCCT GGGAAACCTT TTCTTACAT CTTCAGTGC p39^{st-1}

-617. TAGAAATGT TATGAAACG AAAAGAAT TATAAGGT ATTATAAGG AAACACTCAT TTTCCTCCCA AGAGGCCAA

-537. GATTTCTT TTCTTTTT TTCTCTTCTT TTCTCTTCTT ATTCAAGG AGTATAATTA AATTGCAAGG TAAAAGCTGA

-457. AACGTCTTTT TTATGTTT CTGGAAAGGT CTCTGCCCT GTTGTGATT CCTTCTAGCTT CCACGTTCTT CTATCCAGTT

-377. CGCGACCT TCCCCAGG CCCATTCTT CAAGGCTTCAAGGAGGGT CTCCGGTA AAAGGAAATTC TCAGCACAGA p39^{st-1}

-297. ATCTTCAAC CTCTCTGGAG GCCACCAAG ATCCCTAACG CGGCATGGA GACGAAGCA CTCGGGGGG AGGGAGGCT ACTCACCTCC

-217. GGGGGGGG CCACACCTG SP-1 GGAGAGGGGC GGCACCAAC TGAGGGCGC GGGCTGGGG AGGGAGGCT ACTCACCTCC

-137. CCACCTCCC GGGGGCTACT CAACACGAG SP-1 CACAGGGGGC CACAGGGGGC AGTCTTCTGG CGAGCCTGAGCAGC

-57. [CGCCAGGTGGGAGGTGT CGGTCCGGTCC CGGTCCGGTCC CGGTCCGGTCC CGGTCCGGTCC CGGTCCGGTCC CGGTCCGGTCC

-57. [CGCCAGGTGGGAGGTGT CGGTCCGGTCC CGGTCCGGTCC CGGTCCGGTCC CGGTCCGGTCC CGGTCCGGTCC CGGTCCGGTCC

+23. GGTCGGGGAA CCCCTCGTC CGTGTCTCGTC CGACGGGAC CGGTCCGGTCC CGGTCCGGTCC CGGTCCGGTCC CGGTCCGGTCC

+103. TGTCCCACTC CGGGGACG CCCTCCGG AGTCCCCGG CGGTCCGGTCC CGGTCCGGTCC CGGTCCGGTCC CGGTCCGGTCC

+183. CGGGGGAG GTCCTGGGT TGGGGCTT CCTTCGGGGG AGTCCCCGG CGGTCCGGTCC CGGTCCGGTCC CGGTCCGGTCC

+263. CGATTGGAGC AGAGTGTGG AGCTGGGCTG AGCTGGGCTG AGCTGGGCTG AGCTGGGCTG

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Fig. 2

